academic publishers

INTERNATIONAL JOURNAL OF ARTIFICIAL INTELLIGENCE (ISSN: 2692-5206)

Volume 04, Issue 07, 2024

Published Date: 14-09-2024



"HOW CAN MATHEMATICAL MODELS BE USED TO PREDICT THE PROGRESSION AND IMPACT OF CHRONIC DISEASES IN THE MEDICAL FIELD?"

Mohamed Chahine

Associate professor of the Biological and Chemical Technology, Kursk State Medical University, Kursk, Russian Federation. ORCHID ID: 0000-0001-8504-2418

Nikolaos Tzenios

Professor of Public Health and Medical Research, Charisma University, Grace Bay, Turks and Caicos Islands.

ORCHID ID: 0000-0002-6679-2296

Omasyarifa Binti Jamal Poh

Associate professor of the Obstetrics & Gynecology ICE, Kursk State Medical University, Kursk, Russian Federation. ORCHID ID: 0000-0002-5759-3199

Abstract: Worldwide, healthcare systems face challenges posed by persistent chronic diseases, which have a long-lasting effect on patient health and well-being. Precisely estimating and predicting the origin and impact of chronic diseases is essential for adequate management, funds allocation, and development of policies related to the disease. Mathematical modeling is an important tool for forecasting chronic disease progression, impact, and course. The current article examines several mathematical models, including compartmental, agent-based, and statistical models, emphasizing their distinctive qualities and uses. The models capture intricate relationships between disease-related variables, environmental effects, patient features, and medical therapies. Mathematical models that use data-driven parameter estimates and validation techniques can simulate disease dynamics, forecast patterns for the future, and highlight important variables. Furthermore, these models aid practitioners and other stakeholders in the medical field in decision-making regarding illness prevention, early detection, treatment plans, and allocation of resources. Importantly, mathematical models enhance initiatives' evaluation process and determine their costeffectiveness. The models help reveal underlying mechanisms, advance our understanding of chronic diseases, and enlighten evidence-based medical procedures. The development of mathematical modeling opens up new possibilities for diagnosing and treating chronic diseases. However, data quality, model complexity, and model validation are the weak points of mathematical modeling. Therefore, it is important to discuss future paths and the potential influence of mathematical models on changing how the medical community approaches the management of chronic diseases.

Keywords:

Mathematical Models: Mathematical representations that capture the relationships and dynamics of complex systems, e.g., chronic diseases in the medical field.

Prediction: The use of mathematical models to forecast the future progression and outcomes of chronic diseases based on existing data and assumptions.

Progression refers to how chronic diseases alter and intensify over time and affect patients' health.

Impact: The results and repercussions of chronic illnesses on patients, healthcare systems, and society.

Chronic Ailments: Long-lasting medical problems that require continuing care, such as cancer, diabetes, and respiratory disorders, as well as cardiovascular ailments.

Medical Field: The area of medicine that focuses on healthcare institutions, researchers, and those who diagnose, treat, and manage illnesses.

Compartmental models: Mathematical models divide a population into discrete compartments to describe various disease stages and transitions.

Agent-Based Models: Mathematical models that replicate how individuals operate and communicate in a population, enabling a more in-depth examination of how personality traits and behaviors affect disease development.

Models of system dynamics: Mathematical representations of chains of feedback and interconnections within a system that shed light on the intricate dynamics of chronic diseases and their interactions with other variables.

Parameter estimation: The technique of estimating the parameters in mathematical models based on the information at hand, enabling more precise simulations and predictions.

Validation: The term denotes comparing model results to empirical observations to determine the effectiveness and precision of mathematical models in predicting the course and effects of chronic diseases.

Data-driven: Strategies or techniques that depend upon real-world information to guide the creation, calibration, and validation of mathematical models, ensuring their applicability and accuracy in simulating actual events.

Resource Allocation: The division and distribution of physical, human, and financial assets in the healthcare sector for managing and addressing the burden of chronic diseases.

Interventions: Measures that include lifestyle changes, medication interventions, and public health initiatives. Interventions are actions, regulations, or methods intended to prevent or lessen the progression and effects of chronic diseases.

Interdisciplinary collaborations: The term refers to efforts to build and implement mathematical models in the medical area while utilizing different viewpoints and experiences from professionals in various professions, including medicine, mathematics, statistics, and public health.

Introduction

The persistence of chronic diseases poses challenges for the global medical care systems. Such chronic ailments as cancer, diabetes, cardiovascular illnesses, and respiratory ailments place a significant financial and social burden on patients, healthcare providers, and society. According to Shinde et al. (2020), accurately forecasting the course and effect of the disease is paramount for successful management and treatment. Forecasting enables the medical industry to allocate resources, make knowledgeable decisions, and provide tailored solutions.

In medicine, employing mathematical models to assess and predict how chronic diseases may develop has proven to be an effective method of disease management. Mathematica models incorporate diverse behavioral, biological, and environmental components and offer significant insights into the underlying causes of chronic health disorders (Mohamadou et al., 2020). The models offer important insights into the underlying mechanisms of chronic diseases, aid in identifying important variables affecting the results of the disease, and guide prevention, treatment, and control techniques. Kwok et al. (2019) claim that mathematical models can recreate the disease's natural history by embracing elements such as the disease's length, clinical symptoms, and treatment. The models give medical experts and academics the ability to predict the cause of a disease, gauge the efficacy of various treatment options, and help assess the possible effects of interventions on the development of diseases and population health, such as lifestyle changes or immunization programs. According to Cassidy et al. (2019), mathematical models allow experts to investigate chronic illnesses' economic effects. The models aid in assessing interventions' cost-effectiveness and directing resource distribution by considering factors such as healthcare expenses, decrease in productivity, and overall quality of life. In addition, the models enable evaluation of the long-

term effects of chronic illnesses on people and society, assisting policymakers in creating successful public health plans. Therefore, the development of mathematical modeling will continue to advance our understanding of chronic diseases and aid in creating effective interventions for the population.

Mathematical models provide important information; hence it is significant to understand the applicability and reliability of these models in predicting the impact and progression of chronic diseases. In this regard, comparing and evaluating mathematical models with empirical data and clinical expertise is essential to continuously advance and validate them to guarantee their dependability and trustworthiness. Effective use of mathematical models in the medical field depends on collaboration between physicians, epidemiological researchers, mathematical experts, and policymakers. According to Du & Yuan (2020), the links and interactions between different aspects of a system, such as disease dynamics, individual traits, ecological effects, and treatment alternatives, can be represented mathematically. In the context of chronic diseases, mathematic models help simulate disease progression and make future outcome predictions. Additionally, for the benefit of researchers and doctors, mathematical models allow scenario exploration, evaluation of intervention effectiveness, health resource allocation, and management strategy optimization. This strategy guarantees that the models align with empirical data, clinical knowledge, and developing data, enhancing their value and assisting in more informed decision-making for managing chronic diseases. Therefore, mathematical models in the medical field can be improved by highlighting the necessity for continually validating the results produced and interdisciplinary cooperation.

Advantages of Mathematical Models

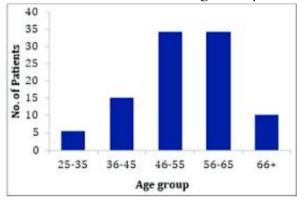
The key benefit of applying mathematical models is forecasting the course and effects of chronic ailments. Mathematical models include various complex and dissimilar data sources, such as clinical data, epidemiological data, and population-level statistics (Brauer et al., 2019). By merging these different sources of information, models can depict the complex interaction of genetic, behavioral, environmental, and healthcare factors that determine the evolution of illness. The accuracy and dependability of forecasts are improved by data integration, which enables a more comprehensive and realistic description of chronic diseases.

Mathematical models have the advantage of simulating and forecasting the development and effects of chronic diseases over a long period. The feature would be particularly important when analyzing chronic problems that gradually worsen over time. With mathematical models, researchers and medical practitioners may imitate symptoms, severity, and prognosis changes as the illness progresses through its many stages (Walters et al., 2018). Mathematical models offer vital insights into the effects of chronic diseases on people, healthcare institutions, and society by forecasting future disease trends. The received information is feasible to organize and carry out efficient healthcare initiatives and plans designed to handle the problems these provide.

To estimate illness prevalence, transmission rates, and the effectiveness of therapies, these models segment the general population into discrete compartments that represent different health conditions and changes. According to Arcede et al. (2020), another benefit of using mathematical models in the medical area is the wide variety of model types available for forecasting the course and effects of chronic diseases. Compartmental models, for example, the well-known Susceptible-Infectious-Recovered (SIR) model used in infectious disease epidemiology, have been established to offer research on non-communicable chronic diseases (Romano et al., 2020). Although, Heydari & Pennock (2018) observed that agent-based models had attracted significant attention, hence increased usage of mathematical models in predicting chronic diseases. These simulations consider such factors as age, sex, genetic predisposition, lifestyle preferences, and treatment adherence and represent the actions and interactions of individual agents within a community. Agent-based models offer a more in-depth examination of disease progression and enable the exploration of customized therapies and targeted tactics by reflecting individual-level variation (Gomez-Vazquez, 2021). By taking into account individual-level determinants and interactions, agent-based models offer a significant advantage in forecasting the development and effects of chronic diseases, enabling a more thorough knowledge of disease dynamics. The said models are customizable and provide uniquely tailored insights, which have a large potential for guiding focused disease management actions and improving patient outcomes in the medical industry.

System dynamics models have also been used to study the intricate dynamics of chronic diseases and their interactions with various causes. These models can be used to examine the long-term consequences of regulations, interventions, and population-level changes because they show the feedback loops and connections present in a system (Darabi & Hosseinichimeh, 2020). Data-driven parameter estimation and validation procedures must be employed to guarantee mathematical models' accuracy and dependability to forecast the course and effects of chronic diseases. These methods are crucial in ensuring that the models successfully capture the dynamics of the diseases and appropriately reflect the patterns of disease progression by comparing the model outputs with actual data. Integrating relevant information from several sources, such as electronic health records, demographic surveys, and illness registries, is crucial to generate accurate parameter estimations. Researchers can compile detailed information on the number of individuals affected by chronic diseases across various age groups by combining data from various sources. The chart below shows the number of individuals per age group affected by chronic diseases.

Chart 1Chronic Diseases in Various Age Groups



Source: Kumari & Bhattacharyya, 2022

According to a study on healthcare systems in Port Blair, Andaman, and Nicobar Islands, individuals aged between 46 and 65 years are highly affected by chronic diseases (Kumari & Bhattacharyya, 2022). Tables 1 and 2 demonstrate the impact of chronic diseases that have become a serious worldwide health concern

Table 1Number of Chronic Diseases Cases Results

Diseases	Dataset name	No. of attributes	Total entries	No. of positive results		Reference
Cancer	Breast Cancer Wisconsin (Diagnostic) Data Set (Dataset 1)	31	569	212	357	(78)
Cancer	Breast Cancer Wisconsin (Dataset 2)	10	699	240	459	(79)
Diabetes	Pima Indians Diabetes Database (Dataset 1)	9	768	268	500	(80)
Diabetes	Diabetes Classification (Dataset 2)	15	390	60	330	(81)
Heart Attack	Heart Disease UCI (Dataset 1)	14	303	138	165	(82)
Heart Attack	Heart Disease Prediction (Dataset 2)	13	270	150	120	(83)
Hepatitis	Hepatitis (Dataset 1)	19	142	80	62	(84)
Hepatitis	Indian Liver Patient Records (Dataset 2)	11	584	168	416	(85)
Kidney	Kidney Disease Dataset (Dataset 1)	26	189	74	115	(86)

Source: Rashid et al., 2022

Table 2

Highest Result of Chronic Cases from Mathematical Artificial Intelligence Algorithms for Constructing Chronic Detection Models

Diseases	Highest results achieved by proposed model
Cancer	98.23%
Diabetes	93.59%
Heart	93.44%
Hepatitis	98.46%
Kidney	98.90%

Source: Rashid et al., 2022

According to the proposed model by Rashid et al. (2022) in Table 1, the models aim to reach the highest level of precision and accuracy in recognizing and managing chronic cases of kidney disease showing a 98.90% highest results (Serebrisky & Wiznia, 2019). Thus, according to a national health survey conducted in 2014, high cholesterol was the chronic condition with the highest prevalence of 58.2%, while emphysema had the lowest prevalence of 4.0% among older adults (see Table 3).

Table 3Chronic Diseases among Older Adults According to CDC National Health Interview Conducted in 2014

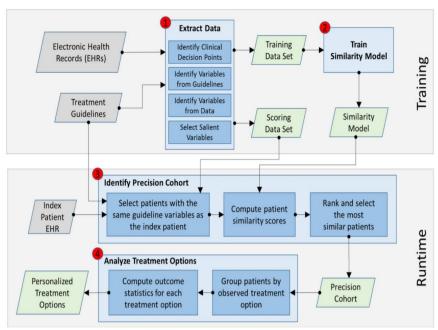
High cholesterol	58.2	
Hypertension	56.7	
Arthritis	48.7	
Cancer	23.1	
Diabetes	20.5	
Heart disease	17.9	
Ulcers	11.3	
Stroke	7.2	
Asthma	6.9	
Kidney disease	5.1	
Chronic bronchitis	5.0	
Emphysema	4.0	

Source: Johnson et al., 2014

Fundamental Concepts of Mathematical Modeling and its Application in Healthcare

Mathematical modeling uses mathematical equations and computer techniques to simulate actual phenomena. The interactions between different elements impacting illness progressions, such as genetic predisposition, environmental exposures, lifestyle decisions, and therapeutic interventions, are captured by mathematical models in healthcare (Ho et al., 2019). The variables and parameters included in these models reflect the traits of the disease, the patient population, and the healthcare system.

Chart 2Personalized Treatment for Chronic Diseases



Source: Ng et al., 2021

Comprehensive understanding of chronic diseases

Mathematical models provide a structure for studying and comprehending the complexities of chronic diseases by considering the complex connections between multiple components. Thus, the models incorporate environmental aspects, genetics, dietary choices, concurrent illnesses, epigenetic factors, and the results of medical therapies. Furthermore, mathematical models facilitate the evaluation of therapeutic approaches and medical interventions (Clarke & Joshu, 2017). The models can predict and evaluate the outcomes of various therapies, aiding in selecting the most effective treatment options by combining information on treatment effectiveness and adherence. Furthermore, the models are effective for healthcare planning and policy formation, as they offer a quantitative framework for evaluating the cost-effectiveness of therapies and allocating resources (Schmitt et al., 2021). Therefore, the ability of mathematical models to represent the complex dynamics and interactions between various components lays the foundation for a thorough understanding of chronic diseases.

Support for decision-making and policy planning

Mathematical models offer useful information that helps healthcare professionals make informed decisions on treatment plans, optimize therapy schedules, and forecast the results of certain therapies. Decision-making for both individual patient treatment and public health planning is greatly aided by mathematical models (Lysaght et al., 2019). Therefore, by combining evidence-based data and applying rigorous mathematical frameworks, the models help medical professionals choose appropriate and individualized treatment regimens that may result in better patient health outcomes.

Furthermore, mathematical models help policymakers in the field of public health allocate resources effectively, develop preventive measures, and assess the cost-effectiveness of various treatments. These models are effective tools for predicting the effects of public health initiatives such as immunization programs or disease prevention plans. By considering numerous scenarios, calculating future results, and determining the most efficient methods to reduce the burden of diseases on the populace, the models assist policymakers in making well-informed decisions (Kretzschmar, 2020). Consequently, the decision-making processes in the healthcare and public health sectors are improved by incorporating mathematical models with evidence-based data. By offering quantitative forecasts and insights, mathematic models lead to more efficient resource allocation, enhanced preventative interventions, and better health outcomes for individuals and communities.

Specific Mathematical Modeling Approaches Commonly Employed in Chronic Disease Prediction

Compartmental models

In mathematical epidemiology, compartmental models are key tools for comprehending disease patterns and forecasting illness progress. The compartmental models divide the population into various compartments based on the states of the diseases, e.g., infected, susceptible, recovered, or deceased (Khailaie et al., 2021). Each of the compartments created represents individuals sharing a common feature or behavior related to the disease. The susceptible compartment reflects individuals at risk of contracting the disease in the context of simulating chronic diseases. The infected compartment represents individuals with the disease, while the recovered compartment represents those who have recovered or are in remission. Therefore, various compartments represent the various stages or states of the disease.

The deterministic compartmental models, which assume changes across compartments and happen at constant rates, are governed by mathematical equations. These models give a streamlined portrayal of disease dynamics and are useful for examining large populations across time. The models enable researchers to calculate disease propagation and important epidemiological variables and forecast future trends (Romano et al., 2020). Additional variables connected to illness-specific aspects, such as disease severity, treatment efficacy, or disease progression rates, can be included in models for chronic diseases.

On the other hand, stochastic compartmental models take into account randomness in the transitions between compartments by recognizing that illness progression might exhibit inherent unpredictability. Stochastic models provide a more accurate depiction of disease dynamics by including probabilistic components like random occurrences and individual-level heterogeneity (Janness et al., 2018). They allow for the study of uncertainty in disease forecasts because they are particularly well-suited to capture the impact of random events and small population numbers. Compartmental models, therefore, offer a framework of mathematics for policy development and choice-making, enabling stakeholders to evaluate the potential efficacy of various disease management and prevention initiatives.

Agent-based models

A more comprehensive individual-level perspective is provided by agent-based models (ABMs), which include heterogeneity and interactions between individuals within a population. Each person (agent) in ABMs has unique characteristics, habits, and disease-related aspects. These simulations of human behavior and interaction over time enable the analysis of complicated scenarios and the detection of emerging illness tendencies (Tracy et al., 2018). ABMs are especially helpful for treating chronic diseases since they may consider things like genetics, social networks, lifestyle choices, and treatment compliance. ABMs can offer insights into customized disease progression, risk stratification, and the impacts of therapies on subpopulations by considering individual-level traits and interactions.

ABMs often concentrate on simulating the behaviors and interactions of single individuals within a population. Compartmental models group populations into uniform groupings, whereas ABMs take into account variability and individual-level dynamics (Asgari-Targhi & Klerman, 2019). Each agent in an ABM represents a distinct entity with particular characteristics, actions, and connections. This individual-level approach makes it possible to simulate complex relationships, social networks, and regional heterogeneity, which greatly impact how chronic diseases develop and affect people. It is crucial not to forget that ABMs also face difficulties with model calibration, parameter estimation, computational complexity, and result interpretation (Asgari-Targhi & Klerman, 2019). ABMs also need a lot of data in order to effectively depict individual-level interactions and behaviors. But despite these difficulties, the distinctive qualities of ABMs make them useful instruments for researching the development and consequences of chronic diseases. They offer a more detailed and nuanced comprehension of disease dynamics, enabling the investigation of intricate relationships and the discovery of tailored therapies and regulations.

System dynamics models

System dynamics models (SDMs) emphasize describing feedback loops and dynamic connections

between variables in intricate structures. SDMs consider various elements in chronic illness modeling, such as biological, behavioral, social, and environmental components that affect disease progression (Liu et al., 2018). SDMs depict these elements as linked parts, enabling the investigation of system behavior over time (Atkins et al., 2018). SDMs allow identifying essential elements and possible remedy locations by modeling feedback loops and nonlinear interactions (Van der Zwet et al., 2022). These models can be utilized for assessing the efficacy of behavior modification programs, forecasting the effects of population-level interventions on the outcomes of chronic diseases, and analyzing the long-term repercussions of policy interventions.

Hybrid models

For certain situations, a mix of modeling techniques may be used to precisely represent the complexity of chronic conditions. Various modeling approaches are combined in hybrid models, like compartmental models with ABMs or SDMs (Liu et al., 2018). This hybridization, incorporating population-level trends and individual-level traits, enables a more thorough knowledge of disease dynamics. For instance, according to Kelleher (2020), a hybrid model may incorporate individual behaviors and interactions through an agent-based component while simulating disease spread at the population level using a compartmental framework. These hybrid models, which consider macro and micro-level dynamics, provide a versatile and effective method for researching chronic diseases.

Diverse Applications of Mathematical Models in Predicting the Progression and Impact of Chronic Diseases

Calculating disease incidence

In order to determine the prevalence of diseases in certain demographic segments, mathematical models are an essential tool. These models make it possible to calculate the prevailing burden of chronic diseases and make projections for future developments by integrating epidemiological data, risk factor profiles, and demographic data (Keogh et al., 2018). These forecasts are crucial for assisting decision-makers and healthcare professionals in comprehending the scope of the problem, effectively allocating resources, and creating focused interventions for high-risk groups. Mathematical models provide healthcare practitioners and decision-makers with information about the scope of the issue by measuring the prevalence of chronic diseases. This data is essential for allocating resources because it identifies regions or populations that need specialized interventions. For instance, healthcare organizations can spend resources to create preventative and treatment methods unique to the requirements of that group if the model predicts a marked rise in the prevalence of a particular chronic disease in a particular demographic group.

Forecasting future healthcare demands

Mathematical models can be used to predict the amount of healthcare that will be required in the future due to chronic conditions cases. These models consider disease prevalence, population growth, demographic aging, and medical breakthroughs. Models can predict the future strain on healthcare systems, including the requirement for hospitalizations, outpatient visits, drugs, and specialized services, by simulating various situations (Ordu et al., 2021). As a result, these estimates aid in planning for healthcare infrastructure, staffing, and budgets.

Optimizing resource allocation strategies

The care and treatment of chronic diseases can benefit from optimizing resource allocation strategies, which mathematical models support. These models consider patient outcomes, healthcare capacity, and treatment costs. Models can uncover solutions that maximize health outcomes while considering restrictions like budgetary constraints by simulating various allocations of resource situations. Models help decide on

the best placement of healthcare workers, medical centers, programs for screening, and prevention efforts.

Simulating "What-if" scenarios

Mathematical models make it possible to simulate "what-if" scenarios, giving us insights into the possible effects of various preventative measures and treatment approaches. Models can investigate the effects of dietary changes, immunization campaigns, early detection initiatives, and focused therapies. Decision-makers can better understand the repercussions and effects of various actions before putting them into practice in real-world contexts by simulating "what-if" scenarios (Freebairn et al., 2018). Decision-makers are able to allocate resources effectively thanks to these simulations' insightful insights about the possible efficacy and cost-effectiveness of initiatives (Freebairn et al., 2018). For example, a mathematical model may mimic the deployment of a certain screening program for a chronic condition and calculate the effect it would have on the rates of early diagnosis, the effectiveness of treatment, and the cost of healthcare. Therefore, models assist decision-makers in choosing the most suitable interventions by assessing the possible advantages, hazards, and cost-effectiveness of alternative tactics through the simulation of these situations.

Importance of reliable data-driven tool

The evolution and effects of chronic diseases can now be predicted using mathematical frameworks in healthcare (Wang et al., 2020). However, these models' precision and dependability depend on data-driven parameter estimation and strict validation procedures. For forecasting and comprehension of the progression and impact of chronic diseases by utilizing mathematical concepts in healthcare, reliable data-driven technologies are of the utmost necessity. These models' reliability and accuracy heavily rely on thorough validation techniques and reliable, data-driven parameter estimates (Wang et al., 2020). To improve the correctness of mathematical models, this article emphasizes the importance of using reliable data sources and carrying out exhaustive calibration and validation procedures. It also highlights the importance of incorporating data from other sources, like demographic surveys and electronic health records (EHRs), to enhance model performance and support well-informed decisions about managing chronic diseases.

For mathematical models to remain precise, it is essential to use correct and trustworthy data sources. The accuracy of parameter estimations is increased using high-quality data that has been rigorously reviewed and acquired from reliable sources, which in turn improves the predictions made by the models. Researchers can better comprehend the complex dynamics of chronic diseases and make more accurate predictions and important insights by depending on complete and current data (Kaplan et al., 2022). The necessary variables and outcomes must be included in the data used for parameter estimation in order for it to appropriately reflect the population being studied.

In addition, precise calibration and validation techniques are just as important as correct data to improve the accuracy and dependability of mathematical models. Model calibration entails altering the model parameters to match the observed data, whereas model validation entails verifying the model's predictions with independent datasets. Calibration can improve models to better represent the intricate dynamics and relationships linked to chronic diseases. Models can also show their ability to generalize beyond the data used for parameter estimation, increasing their applicability and reliability in various scenarios by proving their performance using alternative data sources (Marshall et al., 2018).

Additionally, the effectiveness of mathematical models is greatly enhanced by the inclusion of data from other sources, such as demographic surveys and electronic health records (EHRs). A more in-depth investigation of illness progression and treatment outcomes is possible due to the rich clinical data, longitudinal records, and treatment information provided by EHRs (Yuan et al., 2021). EHRs offer comprehensive patient data, allowing models to consider individual differences and several variables affecting how diseases advance and treatments work. On the other hand, demographic surveys provide useful population-level information that enables researchers to examine illness trends, evaluate the effectiveness of interventions, and create focused disease management plans. Some of the major importance of data-driven tools are discussed below.

Data-driven parameter estimation

The accuracy and dependability of mathematical models for estimating chronic diseases are significantly improved by data-driven parameter estimation. These variables, produced directly from empirical information, are crucial for calibrating and confirming mathematical models. Researchers can enhance the models' prediction power and produce more accurate estimations of the outcomes of chronic diseases by incorporating data-driven factors (Sy et al., 2021). Hence, data-driven parameters are necessary for mathematical models.

Applying statistical techniques like maximum likelihood estimation (MLE) or Bayesian inference is a crucial strategy for data-driven parameter estimation. Using these techniques, researchers can estimate the values of model parameters that best match the observed data (Peng et al., 2020). Whereas Bayesian inference offers a structure for estimating parameters based on beforehand knowledge and updating them using observed data through posterior distributions, MLE, for instance, seeks to find the parameter values that maximize the likelihood of obtaining the observed data (Lio & Liu, 2020). Therefore, statistical techniques are important for data-driven estimations.

These methods reduce the discrepancy between model predictions and actual data to continuously search for optimal model parameter values. The Levenberg-Marquardt method and the genetic algorithm are two examples of optimization algorithms (Dhibi & Amar, 2021). In chronic illness models, accurate parameter estimation depends on the availability and caliber of data. Deriving precise parameters that accurately depict the underlying dynamics of the disease requires comprehensive and representative datasets that capture pertinent factors and outcomes. To ensure that the data used for parameter estimate are reliable and accurate, thorough data collection, preparation, and validation methods are required.

Validation of model predictions

Validation is crucial to guarantee the correctness and dependability of mathematical models' predictions. It entails contrasting model predictions with outside data sources not used for parameter estimation. Researchers can successfully examine a model's capacity to reflect the underlying dynamics of chronic diseases by comparing it to various datasets from various populations, places, or periods (Gao & Yin, 2021). The model's generalizability and structural integrity are improved by this validation procedure, which also reveals any potential restrictions or limitations. Researchers can get insightful knowledge of how the model performs in diverse contexts by extending the range of data utilized for verification, thereby enhancing the model's applicability and reliability. Therefore, the model predictions are validated through this method, which is crucial in proving the validity and usefulness of mathematical models in the study and treatment of chronic diseases.

Reliable data sources for calibrating models

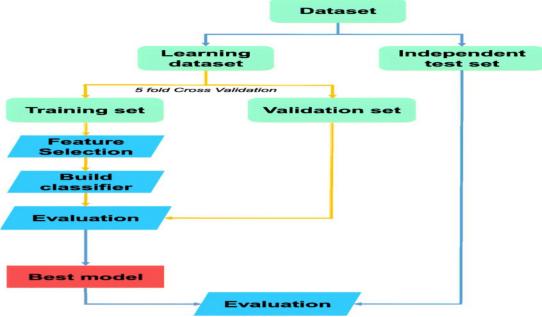
The presence of trustworthy information sources substantially impacts the calibration and validation of mathematical models. Through this mathematical technique, useful sources of patient-level data, including demographics, medical history, treatment records, and results, are electronic health records (EHRs). EHR data integration enables the calibration of disease-specific parameters, individualized risk assessment, and assessment of therapeutic efficacy. Cohort studies and national health surveys are two examples of population studies that offer useful data on illness prevalence, risk factor profiles, and patterns of healthcare consumption (Ali et al., 2018). Therefore, data sources make it possible to calibrate model parameters at the population level and aid in identifying population heterogeneity.

Real-world data integration for improved predictive capabilities

Real-world data improves mathematical models' predictive power and applicability in therapeutic

settings. Combining data from various sources, including wearable technology, surveillance of the environment, social determinants of health, and genomic databases, fills the models with thorough knowledge of disease causation, development, and consequences (Leyens et al., 2017). As a result. Models can make more accurate predictions and help shape specialized strategies for managing chronic diseases by reflecting the intricate relationships between many components. The chart below shows a machine-learning process similar to the mathematical model used to assess the performance of algorithms

Chart 3Machine Learning for Improved Predictability Capabilities of Datasets



Source: Chiesa et al., 2020

Challenges and Limitations Associated with Mathematical Modeling in the Context of Chronic Disease Prediction

Mathematical models have been useful in forecasting the development of chronic disease prediction in medicine. However, it is crucial to understand the difficulties and constraints related to mathematical modeling in this situation (Khan et al., 2019). The following difficulties are addressed.

Model assumptions with uncertainties

Understanding disease dynamics, population behavior, and healthcare systems requires mathematical models. However, because of certain presumptions they function under, which occasionally oversimplify the complicated reality, their projections are subject to risk (Bekiros & Kouloumpou, 2020). The inherent uncertainty in modeling chronic diseases is highlighted using different modeling methodologies and parameter selections that can produce contradictory outcomes. Sensitivity assessments and robustness tests are also essential in assessing the impact of these assumptions on model results to overcome these uncertainties. Promoting transparency in these assessments can help people better grasp the limitations of model assumptions and increase their trust in the results. Therefore, a detailed examination of these uncertainties encourages a more thorough understanding of chronic diseases and assists in making decisions based on the modeling results. Researchers can therefore foster confidence in the results and enable effective communication of the modeling results by noting the existence of uncertainties (Diez-Olivan et al., 2019). Dedication to giving correct information is shown by openly addressing ambiguities in mathematical models, which also fosters a more in-depth comprehension of chronic illness prediction. It enables stakeholders to assess the potential effects of various hypotheses and make well-informed decisions based

Limitations in data availability and quality

For effectively simulating chronic diseases, data accessibility and accuracy are essential. However, there are frequently substantial problems with data availability and quality, which might affect the precision and generalizability of disease prediction models. The fact that data on illness prevalence, risk factors, treatment outcomes, and healthcare utilization are often distorted and biased is one problem. The accuracy and wide applicability of model predictions may be hampered by these data availability limits (Arora et al., 2019). Data are frequently biased for a variety of reasons. A contributing factor is the unequal distribution of healthcare resources and access, which may leave some groups or geographical areas underrepresented or underreported (McCrae, 2018). Models should be checked against actual data, and their assumptions and constraints should be openly disclosed. When using model predictions in decision-making procedures, it is crucial to interpret them in a way that is consistent with clinical and public health viewpoints. Therefore, achieving effective modeling of chronic diseases is significantly hampered by data availability and quality limits. Data biases can result from various factors; hence efforts should be made to enhance data quality measurements. Furthermore, for the efficient application of illness prediction models in decision-making processes, model outputs must be interpreted in the context of clinical and public health perspectives.

Interdisciplinary collaborations

Interdisciplinary Collaborations are necessary to develop accurate and useful mathematical models for chronic disease prediction. Cooperation among epidemiologists, doctors, statisticians, mathematicians, and policymakers is essential to guarantee that models accurately reflect the complex interplay of biological, behavioral, and social aspects (Bekiros & Kouloumpou, 2020). The model's relevance and application are increased by involving domain experts in its creation, parameter estimates, and interpretation. Interdisciplinary partnerships also support the conversion of model predictions into suggestions that healthcare decision-makers can follow.

Model interpretation and communication

It can be difficult to interpret and communicate model results when using mathematical models to predict chronic diseases. Models tend to simplify reality and may not accurately depict all facets of illness patterns. To prevent incorrect interpretation and exploitation of model predictions, stakeholders must be informed about model outputs, uncertainties, and restrictions (Gilbert et al., 2018). Stakeholders can make wise decisions based on model results when assumptions, constraints, and the Context in which models should be utilized are communicated. Different populations and geographical areas should be considered to achieve thorough and representative data gathering. Furthermore, integrating data from various sources and enhancing data exchange among healthcare organizations can assist in reducing data fragmentation and improving the completeness of information. Additionally, the subject of model interpretation is crucial in the context of modeling chronic diseases. Although mathematical models offer useful insights, interpreting them requires careful consideration.

Ethical Considerations and equity

It is crucial to consider equity issues while creating mathematical models to forecast the course and effects of chronic diseases. According to Char, Abramoff, and Feudtner (2020), the varied potential effects of therapies on various specific populations should be explicitly accounted for in these models, along with health inequalities, socioeconomic determinants of health, and other factors. Failure to address these moral and equitable issues may result in ineffective interventions and continuing health disparities. By considering these variables, models can offer insightful information about the distribution of illness burden and aid in developing disparity-mitigation plans. Also, modeling techniques incorporating equality issues can guide

decision-making, resulting in more focused and successful interventions. Regardless of a person's socioeconomic or demographic background, encouraging fairness, social justice, and better health outcomes is impossible without acknowledging the significance of ethics and equity in modeling chronic diseases.

Conclusion

In predicting chronic diseases, mathematical modeling faces many difficulties and constraints, such as ambiguities in model presumptions, restrictions on the quantity and quality of data, and the requirement for interdisciplinary partnerships. Despite these difficulties, mathematical models have a significant potential to advance our comprehension of chronic illnesses and efficient direct interventions. Scholars and medical personnel can use the strength of mathematical models to guide decision-making, optimize resource allocation, and enhance the control of chronic diseases by recognizing these obstacles and actively striving to address them.

In the medical field, mathematical models have generally become effective instruments for forecasting the course and effects of chronic diseases. To offer knowledge about illness progression, medical results, and prospective therapies, these models include a variety of variables, including demographic data, genetic information, environmental variables, and disease-specific parameters. Healthcare practitioners and academics can more fully comprehend the complicated dynamics of chronic diseases, improve treatment plans, and more efficiently allocate resources using mathematical models. Additionally, these models aid in identifying high-risk populations and improve decision-making based on evidence, enabling targeted treatments and preventive actions. Mathematical models have drawbacks and uncertainties, but they have a lot of potential to advance patient care, direct public health initiatives, and eventually lessen the burden of chronic diseases on both people and society.

References:

- 1. Ali, S. A., Soo, C., Agongo, G., Alberts, M., Amenga-Etego, L., Boua, R. P., Choudhury, A., Crowther N. J., Depuur, C., Gómez-Olivé F. X., Guiraud I., Haregu, T. N., Hazelhurst, S., Kahn, K., Khayeka-Wandabwa, C., Kyobutungi, C., Lombard, Z., Mashinya, F., Micklesfield, L., Ramsay, M. (2018). Genomic and environmental risk factors for cardiometabolic diseases in Africa: Methods used for Phase 1 of the AWI-Gen population cross-sectional study. Global Health Action, 11(sup2), 1507133. https://doi.org/10.1080/16549716.2018.1507133
- 2. Arcede, J. P., Caga-Anan, R. L., Mentuda, C. Q., &Mammeri, Y. (2020). Accounting for symptomatic and asymptomatic in a SEIR-type model of COVID-19. Mathematical Modelling of Natural Phenomena, 15, 34. https://doi.org/10.1051/mmnp/2020021
- 3. Arora, P., Boyne, D., Slater, J. J., Gupta, A., Brenner, D. R. & Druzdzel, M. J. (2019). Bayesian networks for risk prediction using real-world data: A tool for precision medicine. Value in Health, 22(4), 439-445. https://doi.org/10.1016/j.jval.2019.01.006
- 4. Asgari-Targhi, A. & Klerman, E. B. (2019). Mathematical modeling of circadian rhythms. Wiley Interdisciplinary Reviews: Systems Biology and Medicine, 11(2), e1439. https://doi.org/10.1002/wsbm.1439
- 5. Atkins, K. E., Lafferty, E. I., Deeny, S. R., Davies, N. G., Robotham, J. V. & Jit, M. (2018). Use of mathematical modelling to assess the impact of vaccines on antibiotic resistance. The Lancet Infectious Diseases, 18(6), e204-e213. https://doi.org/10.1016/S1473-3099(17)30478-4
- 6. Bekiros, S. & Kouloumpou, D. (2020). SBDiEM: A new mathematical model of infectious disease dynamics. Chaos, Solitons & Fractals, 136, 109828. https://doi.org/10.1016/j.chaos.2020.109828
- 7. Brauer, F., Castillo-Chavez, C. & Feng, Z. (2019). Mathematical models in epidemiology. Springer.
- 8. Cassidy, R., Singh, N. S., Schiratti, P. R., Semwanga, A., Binyaruka, P., Sachingongu, N., Chama-Chiliba, C. M., Chalabi, Z., Borghi, J. & Blanchet, K. (2019). Mathematical modelling for health systems research: A systematic review of system dynamics and agent-based models. BMC Health Services Research, 19, 845. https://doi.org/10.1186/s12913-019-4627-7
- 9. Char, D. S., Abràmoff, M. D. & Feudtner, C. (2020). Identifying ethical considerations for machine learning healthcare applications. The American Journal of Bioethics, 20(11), 7-17. https://doi.org/10.1080/15265161.2020.1819469
- 10. Chiesa, M., Maioli, G., Colombo, G. I. & Piacentini, L. (2020). GARS: Genetic algorithm for the

- identification of a robust subset of features in high-dimensional datasets. BMC Bioinformatics, 21, 54. https://doi.org/10.1186/s12859-020-3400-6
- 11. Clarke, M. A. & Joshu, C. E. (2017). Early life exposures and adult cancer risk. Epidemiologic Reviews, 39(1), 11-27. https://doi.org/10.1093/epirev/mxx004
- 12. Darabi, N. & Hosseinichimeh, N. (2020). System dynamics modeling in health and medicine: A systematic literature review. System Dynamics Review, 36(1), 29-73.https://doi.org/10.1002/sdr.1646
- 13. Dhibi, N. & Amar, C. B. (2021). Performance of genetic algorithm and Levenberg-Marquardt method on multi-mother wavelet neural network training for 3D huge meshes deformation: A comparative study. Neural Processing Letters, 53, 2221-2241. https://doi.org/10.1007/s11063-021-10512-y
- 14. Diez-Olivan, A., Del Ser, J., Galar, D. & Sierra, B. (2019). Data fusion and machine learning for industrial prognosis: Trends and perspectives towards Industry 4.0. Information Fusion, 50, 92-111. https://doi.org/10.1016/j.inffus.2018.10.005
- 15. Du, S. Q., & Yuan, W. (2020). Mathematical modeling of interaction between innate and adaptive immune responses in COVID-19 and implications for viral pathogenesis. Journal of Medical Virology, 92(9), 1615-1628. https://doi.org/10.1002/jmv.25866
- 16. Freebairn, L., Atkinson, J. A., Kelly, P. M., McDonnell, G. & Rychetnik, L. (2018). Decision makers' experience of participatory dynamic simulation modelling: Methods for public health policy. BMC Medical Informatics and Decision Making, 18, 131. https://doi.org/10.1186/s12911-018-0707-6
- 17. Gao, Y. & Yin, D. (2021). A full-stage creep model for rocks based on the variable-order fractional calculus. Applied Mathematical Modelling, 95, 435-446. https://doi.org/10.1016/j.apm.2021.02.020
- 18. Gilbert, N., Ahrweiler, P., Barbrook-Johnson, P., Narasimhan, K. P. & Wilkinson, H. (2018). Computational modelling of public policy: Reflections on practice. Journal of Artificial Societies and Social Simulation, 21(1), 14. https://doi.org/10.18564/jasss.3669
- 19. Gomez-Vazquez, J. P. (2021). Agent based modeling to explore intervention strategies for prevention and control of infectious diseases, complex systems modeling in epidemiology [Doctoral dissertation]. University of California.
- 20. Heydari, B. & Pennock, M. J. (2018). Guiding the behavior of sociotechnical systems: The role of agent-based modeling. Systems Engineering, 21(3), 210-226. https://doi.org/10.1002/sys.21435
- 21. Ho, S. M., Lewis, J. D., Mayer, E. A., Bernstein, C. N., Plevy, S. E., Chuang, E., Rappaport, S. M., Croitoru, K., Korzenik, J. R., Krischer, J., Hyams, J. S., Judson, R., Kellis, M., Jerrett, M., Miller, G. W., Grant, M. R., Shtraizent, N., Honig, G., ... Wu, G. D. (2019). Challenges in IBD research: Environmental triggers. Inflammatory bowel diseases, 25(Supplement 2), S13-S23. https://doi.org/10.1093/ibd/izz076
- 22. Johnson, N. B., Hayes, L. D., Brown, K., Hoo, E. C., Ethier, K. A. & Centers for Disease Control and Prevention (CDC) (2014). CDC National Health Report: Leading causes of morbidity and mortality and associated behavioral risk and protective factors—United States, 2005–2013. MMWR Supplements 63(4), 3-27.
- 23. Kelleher, J. (2020). A hybrid agent-based and equation based model for the spread of infectious diseases. JASSS 23(4), 1-25. http://hdl.handle.net/2262/98332
- 24. Keogh, R. H., Szczesniak, R., Taylor-Robinson, D. & Bilton, D. (2018). Up-to-date and projected estimates of survival for people with cystic fibrosis using baseline characteristics: A longitudinal study using UK patient registry data. Journal of Cystic Fibrosis, 17(2), 218-227. https://doi.org/10.1016/j.jcf.2017.11.019
- 25. Khailaie, S., Mitra, T., Bandyopadhyay, A., Schips, M., Mascheroni, P., Vanella, P., Lange, B., Binder, S. C. & Meyer-Hermann, M. (2021). Development of the reproduction number from coronavirus SARS-CoV-2 case data in Germany and implications for political measures. BMC Medicine, 19, 32. https://doi.org/10.1186/s12916-020-01884-4
- 26. Khan, A., Uddin, S. & Srinivasan, U. (2019). Chronic disease prediction using administrative data and graph theory: The case of type 2 diabetes. Expert Systems with Applications, 136, 230-241. https://doi.org/10.1016/j.eswa.2019.05.048
- 27. Kretzschmar, M. (2020). Disease modeling for public health: Added value, challenges, and institutional constraints. Journal of Public Health Policy, 41, 39-51. https://doi.org/10.1057/s41271-019-00206-0

- 28. Kumari, R. & Bhattacharyya, S. (2022). Pandemics and patients with chronic diseases: A study on the health care system in Port Blair, Andaman and Nicobar Islands. Journal of the Anthropological Survey of India, 71(2), 220-235. https://doi.org/10.1177/2277436X221087535
- 29. Kwok, K. O., Tang, A., Wei, V. W., Park, W. H., Yeoh, E. K. & Riley, S. (2019). Epidemic models of contact tracing: Systematic review of transmission studies of severe acute respiratory syndrome and middle east respiratory syndrome. Computational and Structural Biotechnology Journal, 17, 186-194. https://doi.org/10.1016/j.csbj.2019.01.003
- 30. Leyens, L., Reumann, M., Malats, N. & Brand, A. (2017). Use of big data for drug development and for public and personal health and care. Genetic Epidemiology, 41(1), 51-60. https://doi.org/10.1002/gepi.22012
- 31. Lio, W. & Liu, B. (2020). Uncertain maximum likelihood estimation with application to uncertain regression analysis. Soft Computing, 24, 9351-9360. https://doi.org/10.1007/s00500-020-04951-3
- 32. Liu, S., Xue, H., Li, Y., Xu, J. & Wang, Y. (2018). Investigating the diffusion of agent-based modelling and system dynamics modelling in population health and healthcare research. Systems Research and Behavioral Science, 35(2), 203-215. https://doi.org/10.1002/sres.2460
- 33. Lysaght, T., Lim, H. Y., Xafis, V. & Ngiam, K. Y. (2019). AI-assisted decision-making in healthcare: The application of an ethics framework for big data in health and research. Asian Bioethics Review, 11, 299-314. https://doi.org/10.1007/s41649-019-00096-0
- 34. McCrae, R. R. (2018). Method biases in single-source personality assessments. Psychological Assessment, 30(9), 1160. https://doi.org/10.1037/pas0000566
- 35. Mohamadou, Y., Halidou, A. & Kapen, P. T. (2020). A review of mathematical modeling, artificial intelligence and datasets used in the study, prediction and management of COVID-19. Applied Intelligence, 50(11), 3913-3925. https://doi.org/10.1007/s10489-020-01770-9
- 36. Ng, K., Kartoun, U., Stavropoulos, H., Zambrano, J. A. & Tang, P. C. (2021). Personalized treatment options for chronic diseases using precision cohort analytics. Scientific Reports, 11, 1139. https://doi.org/10.1038/s41598-021-80967-5
- 37. Ng, R., Sutradhar, R., Kornas, K., Wodchis, W. P., Sarkar, J., Fransoo, R. & Rosella, L. C. (2020). Development and validation of the chronic disease population risk tool (CDPoRT) to predict incidence of adult chronic disease. JAMA Network Open, 3(6), e204669-e204669. https://doi.org/10.1001/jamanetworkopen.2020.4669
- 38. Ordu, M., Demir, E., Tofallis, C. & Gunal, M. M. (2021). A novel healthcare resource allocation decision support tool: A forecasting-simulation-optimization approach. Journal of the Operational Research Society, 72(3), 485-500. http://dx.doi.org/10.1080/01605682.2019.1700186
- 39. Peng, Y., Fu, M. C., Heidergott, B. & Lam, H. (2020). Maximum likelihood estimation by Monte Carlo simulation: Toward data-driven stochastic modeling. Operations Research, 68(6), 1896-1912. https://doi.org/10.1287/opre.2019.1978
- 40. Rashid, J., Batool, S., Kim, J., Wasif Nisar, M., Hussain, A., Juneja, S. & Kushwaha, R. (2022). An augmented artificial intelligence approach for chronic diseases prediction. Frontiers in Public Health, 10, 860396. https://doi.org/10.3389/fpubh.2022.860396
- 41. Romano, S., Fierro, A. & Liccardo, A. (2020). Beyond the peak: A deterministic compartment model for exploring the Covid-19 evolution in Italy. PLoS One, 15(11), e0241951. https://doi.org/10.1371/journal.pone.0241951
- 42. Schmitt, L., Ochalek, J., Claxton, K., Revill, P., Nkhoma, D. & Woods, B. (2021). Concomitant health benefits package design and research prioritisation: Development of a new approach and an application to Malawi. BMJ Global health, 6(12), e007047. http://dx.doi.org/10.1136/bmjgh-2021-007047
- 43. Shinde, G. R., Kalamkar, A. B., Mahalle, P. N., Dey, N., Chaki, J. & Hassanien, A. E. (2020). Forecasting models for coronavirus disease (COVID-19): A survey of the state-of-the-art. SN Computer Science, 1(197), 1-15. https://doi.org/10.1007/s42979-020-00209-9
- 44. Sy, C., Ching, P. M., San Juan, J. L., Bernardo, E., Miguel, A., Mayol, A. P. Culaba, A., Ubando, A. & Mutuc, J. E. (2021). Systems dynamics modeling of pandemic influenza for strategic policy development: A simulation-based analysis of the COVID-19 case. Process Integration and Optimization for Sustainability, 5, 461-474. https://doi.org/10.1007/s41660-021-00156-9

- 45. Tracy, M., Cerdá, M., & Keyes, K. M. (2018). Agent-based modeling in public health: Current applications and future directions. Annual Review of Public Health, 39, 77-94. https://doi.org/10.1146/annurev-publhealth-040617-014317
- 46. van der Zwet, K., Barros, A. I., van Engers, T. M. & Sloot, P. M. (2022). Emergence of protests during the COVID-19 pandemic: Quantitative models to explore the contributions of societal conditions. Humanities and Social Sciences Communications, 9, 68. https://www.csh.ac.at/publication/emergence-of-protests-during-the-covid-19-pandemic-quantitative-models-to-explore-the-contributions-of-societal-conditions/
- 47. Walters, C. E., Meslé, M. M. & Hall, I. M. (2018). Modelling the global spread of diseases: A review of current practice and capability. Epidemics, 25, 1-8. https://doi.org/10.1016/j.epidem.2018.05.007
- 48. Wang, N., Fu, Y., Zhang, H., & Shi, H. (2020). An evaluation of mathematical models for the outbreak of COVID-19. Precision Clinical Medicine, 3(2), 85-93. https://doi.org/10.1093/pcmedi/pbaa016